

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/816,124

CRF Processing Date: 3/12/2002

Edited by: [Signature]

Verified by: [Signature] (STIC staff) 47

☐

Changed a file from non-ASCII to ASCII

☐

Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐

Edited a format error in the Current Application Data section, specifically:

☐

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____

☐

Added the mandatory heading and subheadings for "Current Application Data".

☐

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

☐

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

☐

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

☐

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐

Inserted colons after headings/subheadings. Headings edited included:

☐

Deleted extra, invalid, headings used by an applicant, specifically:

☒

Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____

☐

Inserted mandatory headings, specifically: _____

☐

Corrected an obvious error in the response, specifically: _____

☐

Edited identifiers where upper case is used but lower case is required, or vice versa.

☐

Corrected an error in the Number of Sequences field, specifically: _____

☐

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐

Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐

Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

1600 1645
3/12/2002
RECEIVED
MAR 15 2002
TECH CENTER 1600/2900

ENTERED



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/816,124

DATE: 03/12/2002

TIME: 17:25:55

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\03122002\I816124.raw

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3 <110> APPLICANT: Nagasawa, Yasuo
4     Yoshida, Hideaki
6 <120> TITLE OF INVENTION: Method for detecting and isolating genes
8 <130> FILE REFERENCE: S1-801PCT
10 <140> CURRENT APPLICATION NUMBER: US 09/816,124
11 <141> CURRENT FILING DATE: 2001-03-26
13 <150> PRIOR APPLICATION NUMBER: PCT/JP97/04126
14 <151> PRIOR FILING DATE: 1997-11-12
16 <150> PRIOR APPLICATION NUMBER: JP 1996-305163
17 <151> PRIOR FILING DATE: 1996-11-15
19 <160> NUMBER OF SEQ ID NOS: 4
21 <170> SOFTWARE: PatentIn version 3.0
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 29
25 <212> TYPE: DNA
26 <213> ORGANISM: Artificial Sequence
28 <220> FEATURE:
29 <223> OTHER INFORMATION: Artificially synthesized primer sequence
31 <400> SEQUENCE: 1
32 ataagctttt cacatgagcg aaaaataca                29
34 <210> SEQ ID NO: 2
35 <211> LENGTH: 29
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial Sequence
39 <220> FEATURE:
40 <223> OTHER INFORMATION: Artificially synthesized primer sequence
42 <400> SEQUENCE: 2
43 atggatccct attgtaaccc gctgaagt                29
45 <210> SEQ ID NO: 3
46 <211> LENGTH: 38
47 <212> TYPE: DNA
48 <213> ORGANISM: Artificial Sequence
50 <220> FEATURE:
51 <223> OTHER INFORMATION: Artificially synthesized primer sequence
53 <400> SEQUENCE: 3
54 atgtctcgag aattcagtaa cccaggcatt attttatc        38
56 <210> SEQ ID NO: 4
57 <211> LENGTH: 30
58 <212> TYPE: DNA
59 <213> ORGANISM: Artificial Sequence
61 <220> FEATURE:
62 <223> OTHER INFORMATION: Artificially synthesized primer sequence
64 <400> SEQUENCE: 4

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RAW SEQUENCE LISTING

DATE: 03/12/2002

PATENT APPLICATION: US/09/816,124

TIME: 17:25:55

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\03122002\I816124.raw

65 ttgtcctaga agcttgtgtg ctctgctgtc

30

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/816,124

DATE: 03/12/2002

TIME: 17:25:56

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\03122002\I816124.raw



1645

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/816,124

DATE: 03/01/2002

TIME: 15:07:13

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\03012002\I816124.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Nagasawa, Yasuo
 4 Yoshida, Hideaki
 6 <120> TITLE OF INVENTION: Method for detecting and isolating genes
 8 <130> FILE REFERENCE: S1-801PCT
 10 <140> CURRENT APPLICATION NUMBER: US 09/816,124
 11 <141> CURRENT FILING DATE: 2001-03-26
 13 <150> PRIOR APPLICATION NUMBER: PCT/JP97/04126
 14 <151> PRIOR FILING DATE: 1997-11-12
 16 <150> PRIOR APPLICATION NUMBER: JP 1996-305163
 17 <151> PRIOR FILING DATE: 1996-11-15
 19 <160> NUMBER OF SEQ ID NOS: 4
 21 <170> SOFTWARE: PatentIn version 3.0

ERRORED SEQUENCES

56 <210> SEQ ID NO: 4
 57 <211> LENGTH: 30
 58 <212> TYPE: DNA
 59 <213> ORGANISM: Artificial Sequence
 61 <220> FEATURE:
 62 <223> OTHER INFORMATION: Artificially synthesized primer sequence
 64 <400> SEQUENCE: 4
 65 ttgtcctaga agcttggtg ctctgctgtc 30

E--> 67 38
 E--> 69 1/2

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/816,124

DATE: 03/01/2002

TIME: 15:07:14

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\03012002\I816124.raw

L:67 M:254 E: No. of Bases conflict, LENGTH:Input:38 Counted:30 SEQ:4

M:254 Repeated in SeqNo=4

L:69 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2

L:69 M:252 E: No. of Seq. differs, <211>LENGTH:Input:30 Found:31 SEQ:4